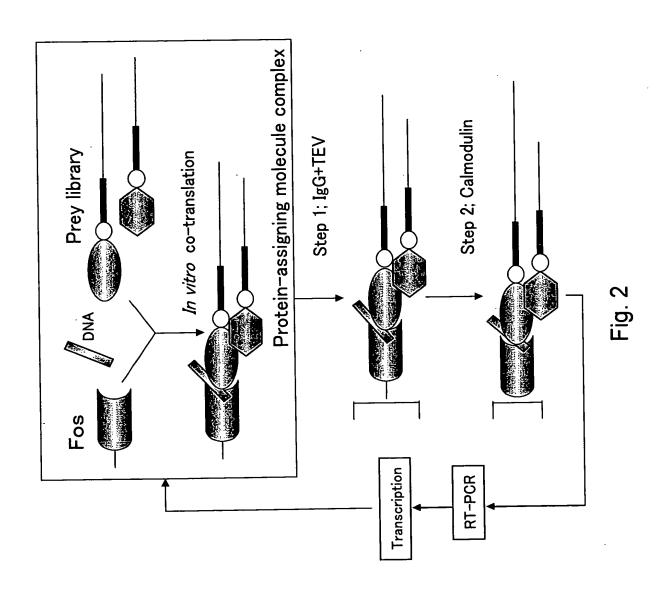
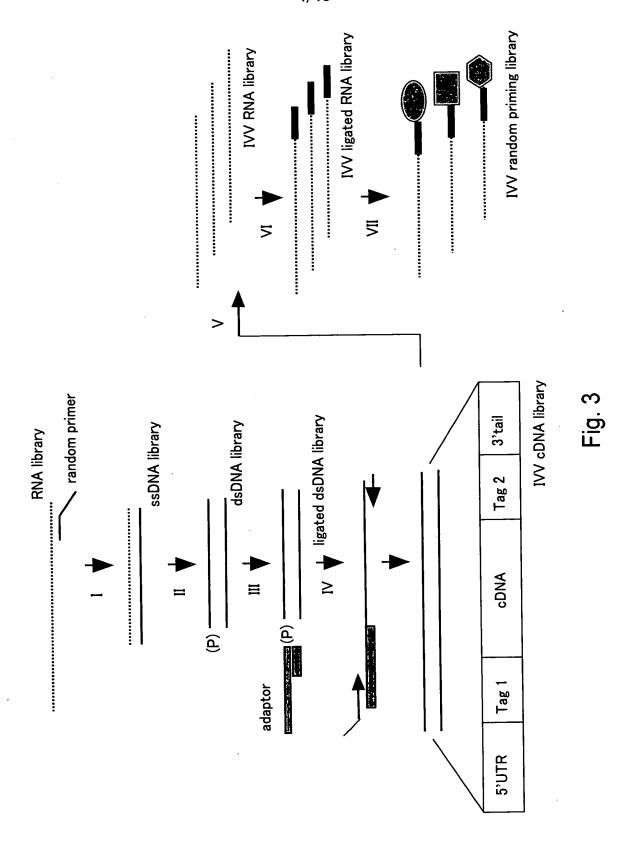
Amino acid SEQ ID NO.	Name of protein/gene, accession No.	leu zipper	Nucleic acid SEQ ID NO.	Num- ber of clones	Alternate Symbols & Ailas
1~14	Mus musculus fip-cx	0	23(1), 24(2–1), 25(2–2), 26(2–3), 27(3), 28(4), 29(5), 30(6), 31(7), 32(8), 33(9), 34(10), 35(11), 36(12), 37(13), 38(14)	59	Frame shift (mage-d3, mRNA, AF319977. melanoma antigen, family D, 3-like, AK047777. trophinin, NM_019548. Trol, Maged3, Maged3I magphinin-alpha, mRNA, AF241245. magphinin-beta2 mRNA, AF288605. magphinin-gamma mRNA, AF288606. trophinin-2 mRNA)
15~19	Mus musculus eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d), mRNA, NM 023240.	0	39(15), 40(16), 41(17), 42(18), 43(19)	rc	5730529A16Rik
20~22	Mus musculus schwannomin interacting protein 1 (Schip1), mRNA, NM_013928.	0	44(20), 45(21), 46(22)	ო	Nf2ip, SCHIP-1
47~56	Mus musculus fip-cx.1	0	104(47), 105(48), 106(49), 107(50-1), 108(50-2), 109(50-3), 110(50-4), 111(50- 5), 112(51-1), 113(51-2), 114(52), 115(53), 116(54), 117(55), 118(56)	15	Frame shift (mage-d3, mRNA, AF319977. melanoma antigen, family D, 3-like, AK047777. trophinin, NM_019548. Trol, Maged3, Maged31)
57~76	Mus musculus fip-cx.2	0	119(57), 120(58), 121(59), 122(60), 123(61), 124(62–1), 125(62–2), 126(63), 127(64), 128(65), 129(66), 130(67), 131(68), 132(69), 133(70), 134(71), 135(72), 136(73), 137(74–1), 138(74–2), 139(75), 140(76)	31	Frame shift (magphinin-alpha, mRNA, AF241245. magphinin mRNA AB032477. magphinin-beta2 mRNA, AF288605. magphinin-gamma mRNA, AF288606. trophinin-2 mRNA)

Fig. 1A

Name of protein/gene, accession No.	No.	Leu zipper	Nucleic acid SEQ ID NO.	Num- ber of clones	Alternate Symbols & Ailas
Mus musculus optineurin (Optn), NM_181848			141(77), 142(78), 143(79), 144(80). 145(81)	9	NRP, FIPZ, HYPL, 4930441007KiK, TFIIIA-INTP
Mus musculus similar to small nuclear RNA activating complex, polypeptide 5, 19kDa; Small nuclear RNA activating complex, polypeptide 5, XM_284503.1		14	146(82), 147(83), 148(84)	2	Snapc5, 2010103A03Rik
4Rik , BC026483 O		7	149(85), 150(86)	-	MGC31554
Rattus norvegicus similar to hypothetical O 15 protein FLJ32000, XM_342896.1		=	151(87), 152(88), 153(89)	2	
CAAX 2 ×		≝_	154(90), 155(91)	-	Rit2
Mus musculus isolate 1 cytochrome b gene, O 11 partial , mitocondorial gene, AF540912.1		=	156(92), 157(93)	-	
×		=	158(94), 159(95)	-	Apoe
s amyloid beta (A4) precursor x)5490.1		Ξ	160(96), 161(97)	-	Adap, Cvap, Abeta, appican, betaAPP, protease nexin II
Mus musculus DnaJ homolog, subfamily A, x member 2, BC003420	-	16	162(98), 163(99)	-	Hsp40 homolog, subfamily A, member 2, DNAJ, DNJ3, mDj3, Dnaj3, HIRIP4, PRO3015, DNA J protein
MUs musculus fip-c10, XM_136911 x 16		=	164(100), 165(101)	-	Mus musculus similar to KIAA1209 protein
Mus musculus fip-c4 × 16		=_	166(102)	-	Genome (Mouse DNA sequence from clone RP23-185C16 on chromosome 4)
Mus musculus fip-c18 ×			167(103)	-	Genome (Mus musculus chromosome 18, clone RP24-572G3, AC102422.10)

Fig. 1B





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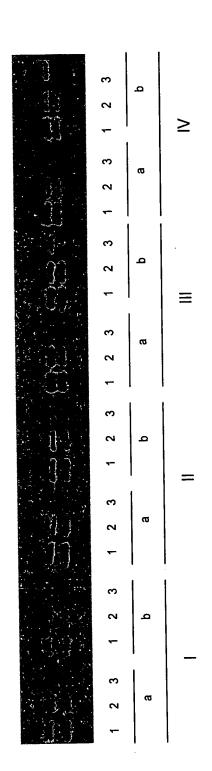
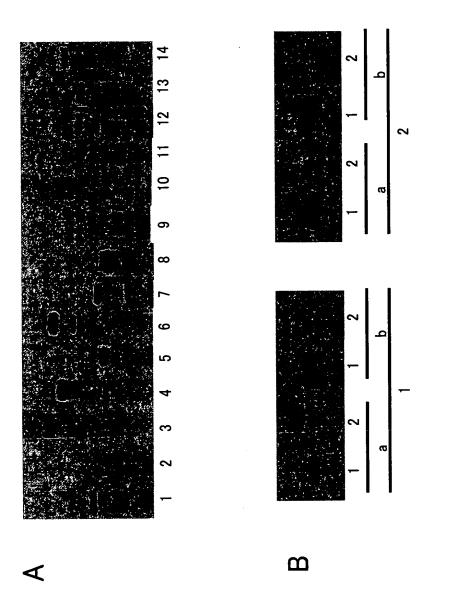


Fig. 4

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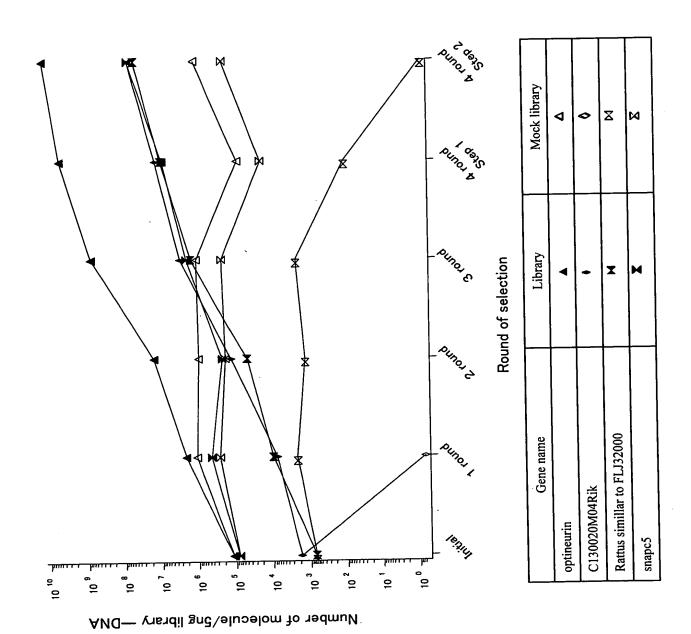
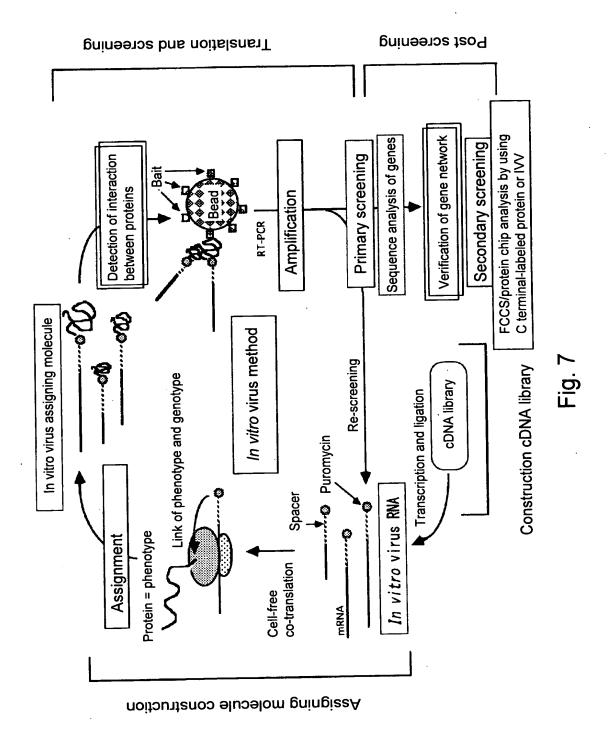


Fig. 6



A Translation template

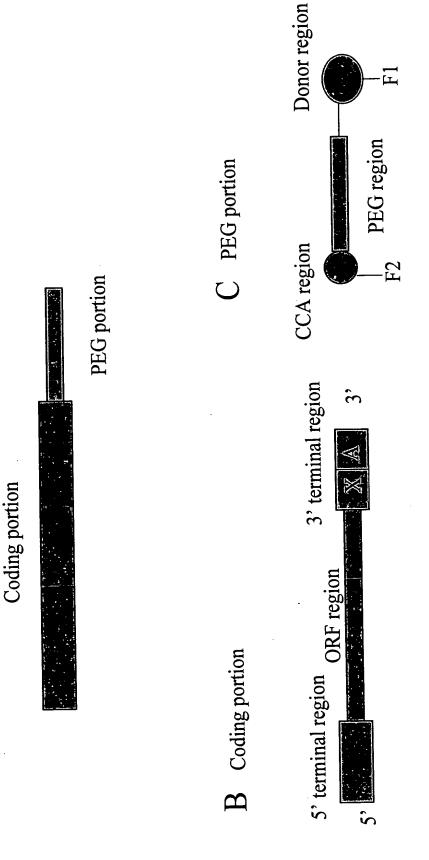


Fig. 8

A Protein of which C terminal is modified with translation template

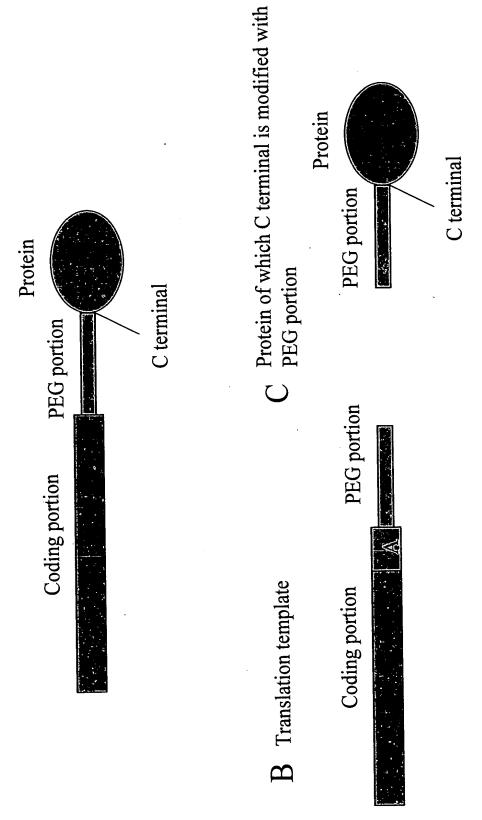


Fig. 9

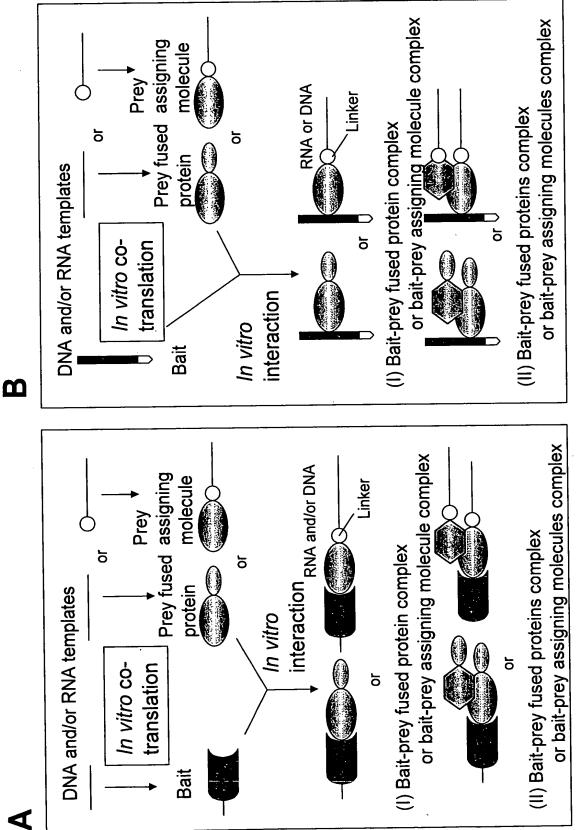


Fig. 10

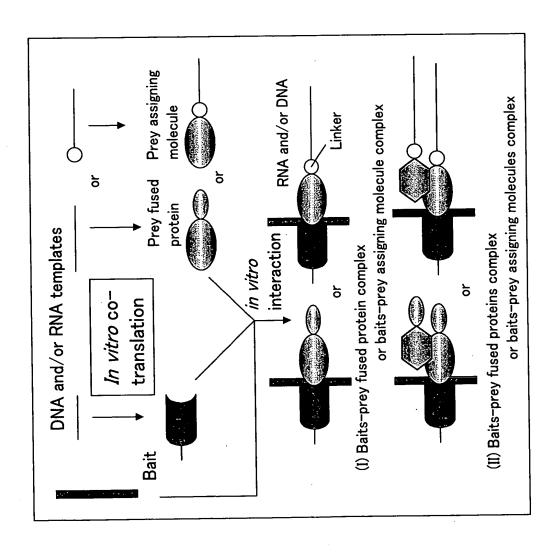


Fig. 11

